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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/910,059

DATE: 10/29/2001
TIME: 15:18:02

Input Set : A:\pto_vsk.txt
Output Set: N:\CRF3\10292001\I910059.raw

3 <110> APPLICANT: Copley, Clive G
4 Edge, Michael Derek
5 Emery, Stephen Charles
7 <120> TITLE OF INVENTION: Monoclonal Antibody to CEA, Conjugates Comprising Said Antibody, and
8 Their Therapeutic use in an Adept System
10 <130> FILE REFERENCE: 1991-209
12 <140> CURRENT APPLICATION NUMBER: US 09/910,059
13 <141> CURRENT FILING DATE: 2001-07-23
15 <150> PRIOR APPLICATION NUMBER: US 09/171,945
16 <151> PRIOR FILING DATE: 1998-10-29
18 <150> PRIOR APPLICATION NUMBER: PCT/GB97/01165
19 <151> PRIOR FILING DATE: 1997-04-29
21 <150> PRIOR APPLICATION NUMBER: GB 9703103.3
22 <151> PRIOR FILING DATE: 1997-02-14
24 <150> PRIOR APPLICATION NUMBER: GB9609405.7
25 <151> PRIOR FILING DATE: 1996-05-04
27 <160> NUMBER OF SEQ ID NOS: 131
29 <170> SOFTWARE: PatentIn version 3.1
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32 <211> LENGTH: 32
33 <212> TYPE: DNA
34 <213> ORGANISM: Artificial Sequence ✓
36 <220> FEATURE:
37 <223> OTHER INFORMATION: light chain cDNA foward primer ✓
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44 <211> LENGTH: 31
45 <212> TYPE: DNA
46 <213> ORGANISM: Artificial Sequence ✓
48 <220> FEATURE:
49 <223> OTHER INFORMATION: heavy chain cDNA foward primer ✓
51 <400> SEQUENCE: 2
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56 <211> LENGTH: 34
57 <212> TYPE: PRT
58 <213> ORGANISM: Mus musculus
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63 1 5 10 15
66 Ser Val Lys Leu Ser Cys Thr Ala Ser Gly Phe Asn Ile Lys Asp Asn
67 20 25 30
70 Tyr Met
74 <210> SEQ ID NO: 4
75 <211> LENGTH: 24
76 <212> TYPE: DNA

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77 <213> ORGANISM: Artificial Sequence
79 <220> FEATURE:
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89 <213> ORGANISM: Artificial Sequence
91 <220> FEATURE:
92 <223> OTHER INFORMATION: light chain cDNA backward primer
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100 <212> TYPE: DNA
101 <213> ORGANISM: Artificial Sequence
103 <220> FEATURE:
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107 aggttsmarct gcagsagtcw gg 22
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112 <212> TYPE: DNA
113 <213> ORGANISM: Artificial Sequence
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122 <210> SEQ ID NO: 8
123 <211> LENGTH: 357
124 <212> TYPE: DNA
125 <213> ORGANISM: Mus musculus
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130 ataacctgca gtgccagtc aagtgttaact tacatgcact ggttccagca gaagccaggc 120
132 acttctccca aactctggat ttatagcaca tccaaacctgg cttctggagt ccctgctcgc 180
134 ttcaatgtggca gtggatctgg gacctcttac tctctcacaa tcagccgaat ggaggctgaa 240
136 gatgctgcca cttattactg ccagcaaagg agtacttacc cgctcacgtt cggtgctggg 300
138 accaagctgg agctgaaaacg ggctgatgct gcaccaactg tatccatott caagctt 357
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142 <211> LENGTH: 108
143 <212> TYPE: PRT
144 <213> ORGANISM: Mus musculus
146 <400> SEQUENCE: 9
148 Asp Ile Glu Leu Thr Gln Ser Pro Ala Ile Met Ser Ala Ser Pro Gly
149 1 5 10 15
152 Glu Lys Val Thr Ile Thr Cys Ser Ala Ser Ser Ser Val Thr Tyr Met
153 20 25 30

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156 His Trp Phe Gln Gln Lys Pro Gly Thr Ser Pro Lys Leu Trp Ile Tyr
157 35 40 45
160 Ser Thr Ser Asn Leu Ala Ser Gly Val Pro Ala Arg Phe Ser Gly Ser
161 50 55 60
164 Gly Ser Gly Thr Ser Tyr Ser Leu Thr Ile Ser Arg Met Glu Ala Glu
165 65 70 75 80
168 Asp Ala Ala Thr Tyr Tyr Cys Gln Gln Arg Ser Thr Tyr Pro Leu Thr
169 85 90 95
172 Phe Gly Ala Gly Thr Lys Leu Glu Leu Lys Arg Ala
173 100 105
176 <210> SEQ ID NO: 10
177 <211> LENGTH: 360
178 <212> TYPE: DNA
179 <213> ORGANISM: Mus musculus
181 <400> SEQUENCE: 10
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184 tcctgcacag cttctggctt caacattaaa gacaactata tgcactgggt gaaggcagg 120
186 cctgaacagg gcctggagtg gattgcattt attgatcctg agaatggta tactgaatat 180
188 gccccgaagt tccggggcaa ggccactttt actgcagact catcctccaa cacagcctac 240
190 ctgcaccta gcagcctgac atctgaggac actgcgtctt attactgtca tgtcctgatc 300
192 tatgctggtt atttggctat ggactactgg ggtcaaggaa cctcagtgcg cgttcctca 360
195 <210> SEQ ID NO: 11
196 <211> LENGTH: 120
197 <212> TYPE: PRT
198 <213> ORGANISM: Mus musculus
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203 1 5 10 15
206 Ser Val Lys Leu Ser Cys Thr Ala Ser Gly Phe Asn Ile Lys Asp Asn
207 20 25 30
210 Tyr Met His Trp Val Lys Gln Arg Pro Glu Gln Gly Leu Glu Trp Ile
211 35 40 45
214 Ala Trp Ile Asp Pro Glu Asn Gly Asp Thr Glu Tyr Ala Pro Lys Phe
215 50 55 60
218 Arg Gly Lys Ala Thr Leu Thr Ala Asp Ser Ser Ser Asn Thr Ala Tyr
219 65 70 75 80
222 Leu His Leu Ser Ser Leu Thr Ser Glu Asp Thr Ala Val Tyr Tyr Cys
223 85 90 95
226 His Val Leu Ile Tyr Ala Gly Tyr Leu Ala Met Asp Tyr Trp Gly Gln
227 100 105 110
230 Gly Thr Ser Val Ala Val Ser Ser
231 115 120
234 <210> SEQ ID NO: 12
235 <211> LENGTH: 39
236 <212> TYPE: DNA
237 <213> ORGANISM: Artificial Sequence
239 <220> FEATURE:
240 <223> OTHER INFORMATION: light chain primer
242 <400> SEQUENCE: 12

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Input Set : A:\pto_vsk.txt
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243 aagctttccc gcggggacat tgagctcacc cagtctcca	39
246 <210> SEQ ID NO: 13	
247 <211> LENGTH: 30	
248 <212> TYPE: DNA	
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251 <220> FEATURE:	
252 <223> OTHER INFORMATION: light chain primer ✓	
254 <400> SEQUENCE: 13	
255 aagcttctcg agcttggtcc cagcaccgaa	30
258 <210> SEQ ID NO: 14	
259 <211> LENGTH: 36	
260 <212> TYPE: DNA	
261 <213> ORGANISM: Artificial Sequence ✓	
263 <220> FEATURE:	
264 <223> OTHER INFORMATION: heavy chain primer ✓	
266 <400> SEQUENCE: 14	
267 aagcttggaa ttcaagtgtga ggtgcagctg cagcag	36
270 <210> SEQ ID NO: 15	
271 <211> LENGTH: 33	
272 <212> TYPE: DNA	
273 <213> ORGANISM: Artificial Sequence ✓	
275 <220> FEATURE:	
276 <223> OTHER INFORMATION: heavy chain primer ✓	
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279 aagcttcgag ctcacggcga ctgagggttcc ttg	33
282 <210> SEQ ID NO: 16	
283 <211> LENGTH: 705	
284 <212> TYPE: DNA	
285 <213> ORGANISM: Artificial Sequence ✓	
287 <220> FEATURE:	
288 <223> OTHER INFORMATION: chimaeric light chain sequence ✓	
290 <400> SEQUENCE: 16	
291 atggattttc aagtgcagat ttctcagttc ctgctaatca gtgcttcagt cataatgtcc	60
293 cgcggggaca tttagctcac ccagtctcca gcaatcatgt ctgcacatctcc aggggagaag	120
295 gtcaccataa cctgcagtgc cagctcaagt gtaacttaca tgcactgggtt ccagcagaag	180
297 ccaggcactt ctcccaaact ctggatttat agcacatcca acctggcttc tggagtcct	240
299 gctcgcttca gtggcagtgg atctgggacc tcttactctc tcacaatcag ccgaatggag	300
301 gctgaagatg ctgccactta ttactgccag caaaggagta cttacccgct cacgttcgg	360
303 gctgggacca agctcgagat caaaacggact gtggctgcac catctgtt catcttcccg	420
305 ccatctgtat agcagttgaa atctggaaact gcctctgtt tgcctgtt gaataacttc	480
307 tatcccagag aggccaaagt acagtggaaag gtggataacg ccctccaatc gggtaactcc	540
309 caggagatg tcacagagca ggacagcaag gacagcacct acagcctcag cagcaccctg	600
311 acgctgagca aagcagacta cgagaaacac aaagtctacg cctgcgaagt caccatcag	660
313 ggctcgagtt cggccgtcac aaagagcttc aacaggggag agtgt	705
316 <210> SEQ ID NO: 17	
317 <211> LENGTH: 235	
318 <212> TYPE: PRT	
319 <213> ORGANISM: Artificial Sequence	
321 <220> FEATURE:	

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Input Set : A:\pto_vsk.txt
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322 <223> OTHER INFORMATION: chimaeric light chain sequence
 324 <400> SEQUENCE: 17
 326 Met Asp Phe Gln Val Gln Ile Phe Ser Phe Leu Leu Ile Ser Ala Ser
 327 1 5 10 15
 330 Val Ile Met Ser Arg Gly Asp Ile Glu Leu Thr Gln Ser Pro Ala Ile
 331 20 25 30
 334 Met Ser Ala Ser Pro Gly Glu Lys Val Thr Ile Thr Cys Ser Ala Ser
 335 35 40 45
 338 Ser Ser Val Thr Tyr Met His Trp Phe Gln Gln Lys Pro Gly Thr Ser
 339 50 55 60
 342 Pro Lys Leu Trp Ile Tyr Ser Thr Ser Asn Leu Ala Ser Gly Val Pro
 343 65 70 75 80
 346 Ala Arg Phe Ser Gly Ser Gly Ser Gly Thr Ser Tyr Ser Leu Thr Ile
 347 85 90 95
 350 Ser Arg Met Glu Ala Glu Asp Ala Ala Thr Tyr Tyr Cys Gln Gln Arg
 351 100 105 110
 354 Ser Thr Tyr Pro Leu Thr Phe Gly Ala Gly Thr Lys Leu Glu Ile Lys
 355 115 120 125
 358 Arg Thr Val Ala Ala Pro Ser Val Phe Ile Phe Pro Pro Ser Asp Glu
 359 130 135 140
 362 Gln Leu Lys Ser Gly Thr Ala Ser Val Val Cys Leu Leu Asn Asn Phe
 363 145 150 155 160
 366 Tyr Pro Arg Glu Ala Lys Val Gln Trp Lys Val Asp Asn Ala Leu Gln
 367 165 170 175
 370 Ser Gly Asn Ser Gln Glu Ser Val Thr Glu Gln Asp Ser Lys Asp Ser
 371 180 185 190
 374 Thr Tyr Ser Leu Ser Ser Thr Leu Thr Leu Ser Lys Ala Asp Tyr Glu
 375 195 200 205
 378 Lys His Lys Val Tyr Ala Cys Glu Val Thr His Gln Gly Leu Ser Ser
 379 210 215 220
 382 Pro Val Thr Lys Ser Phe Asn Arg Gly Glu Cys
 383 225 230 235

386 <210> SEQ ID NO: 18

387 <211> LENGTH: 765

388 <212> TYPE: DNA

389 <213> ORGANISM: Artificial Sequence

391 <220> FEATURE:

392 <223> OTHER INFORMATION: chimaeric HuIgG2 Fd construct

394 <400> SEQUENCE: 18

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397 gtgcagctgc agcartcagg ggcagagctt	gtgagggtcag gggcctcagt	caagttgtcc	120
399 tgcacagctt ctggcttcaa cattaaagac	aactatatgc actgggtgaa	gcagaggcct	180
401 gaacaggccc tggagtgat tgcatggatt	gatcctgaga atggtgatac	tgaatatgcc	240
403 ccgaagttcc gggcaaggc cacttgact	gcagactcat cctccaacac	agccatcac	300
405 cacctcagca gcctgacatc tgaggacact	gccgtctatt actgtcatgt	cctgatctat	360
407 gctggttatt tggctatgga ctactgggt	caaggaacct cagtcgcgt	gagtcggct	420
409 agcaccaagg gaccatcggt ctccccctg	gccccctgct ccaggagcac	ctccgagagc	480
411 acagccgccc tgggctgcct ggtcaaggac	tacttccccg aaccggtgac	ggtgtcgtgg	540
413 aactcaggcg ctctgaccag cggcgtgcac	accttcccgg ctgtcctaca	gtcctcagga	600

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/910,059

DATE: 10/29/2001
TIME: 15:18:03

Input Set : A:\pto_vsk.txt
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